

WB anti P-FKHR

16

Figure 2A

18

WB anti P-FKHR

19 20

Figure 2B

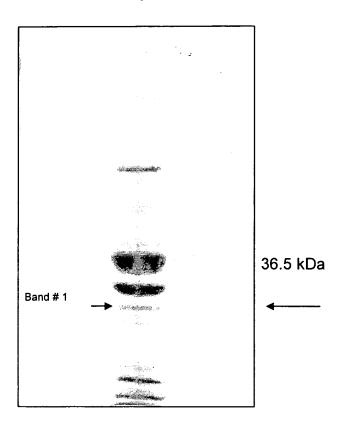


Figure 2C

Q9Y365 FKHR1

QHADSLENIDESAVAESR [SEQ ID NO:2]

RRAASMDNNSKFAKSRSR

[SEQ ID NO:3]

Identity: 6 (33 %)

Strongly similar: 6 (33 %) Weakly similar: 2 (11%)

Different: 4 (22 %)

Figure 2D

DVGYYS WRCPKPLKNRDVITLRS WLPMGADYIIMNYSVKHPKYPPRKDLVRAVSIQTGYLIQ MSTRAKKLRRI WRILEEEESVAGAVQTLLLRSQEGGVTSAAASTLSEPPRRTQESRTRTRA STGPKSCVITYLAQVDPKGSLPKWVVNKSSQFLAPKAMKKMYKACLKYPEWKQKHLPHFK VSVWVQAVEMDRTLHKIKCRMECCDVPAETLYDVLHDIEYRKKWDSNVIETFDIARLTVNA LGLPTLPM68 EKLAASTEPQGPRPVLGRESVQVPDDQDFRSFRSECEAEVG WNLTYSRAG PWLHPEQSPLPSLALSELSVQHADSLENIDESAVAESREERMGGAGGEGSDDDTSLT [SEQ ID NO:6]

Figure 3A

293T HA-Fors
SKBR3

WB anti P-FKHR

15.

Figure 3B

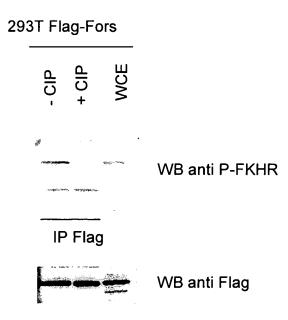


Figure 3C

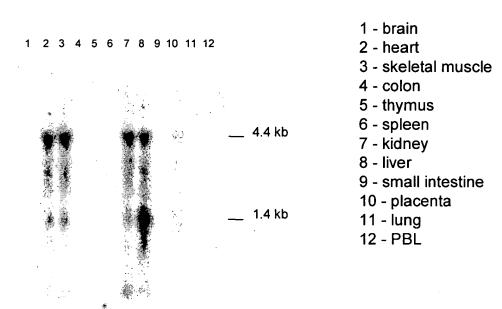
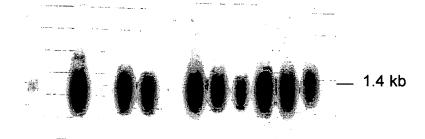
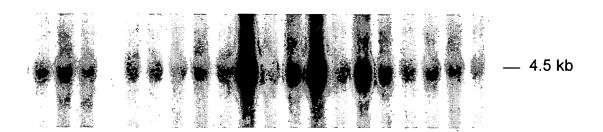


Figure 3D





Fors1



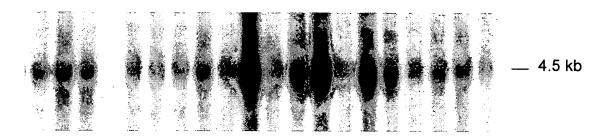
ErbB2

Figure 4A





Fors1



ErbB2

Figure 4B

MEKLAASTEPQG PRPVLGRESVQVPDDQDFRSFRSECEAEVG WNLTYSRAGVSV WVQAVE -LOOPALAGAD WOLLVETSGISIY-RLLD MELAAGSFSEEQFWEACAE-Fors1

KKTGLHEYKVFGVLEDCSPTLLADIYMDSDYRKQWDQYVKELYE--QECNGETVVYWEVK MDRTLHKIKCRMECCDVPAETLYDVLHDIEYRKKWDSNVIETFDIARLTVNADVGYYSWR ************ Fors1 Pctp

YPFPMSNRDYVYLRQRRDLDMEGRKHWILARSTSMPQLGERSGVIRVKQYKQSLAIESD CPKPLKNRDVITLRSWLPMGAD---YIIMNYSVKHPKYPPRKDLVRAVSIQTGYLLOST Fors1

GPKSCVITYLAQVDPKGSLPKWVVNKSSQFLAPKAMKKMYKACLKYPEWKQKHLPHFKPW GKKGSKVFMYYFDNPGGQIPS WLINWAAKNGVPNFLKDMARACQNYLKKT--*** *** ** ** ** ** *** * *** Pctp

LHPEQSPLPSLALSELSVQHADSLENIDESAVAESREERMGAGGGEGSDDDTSLT

Fors1

Figure 5A

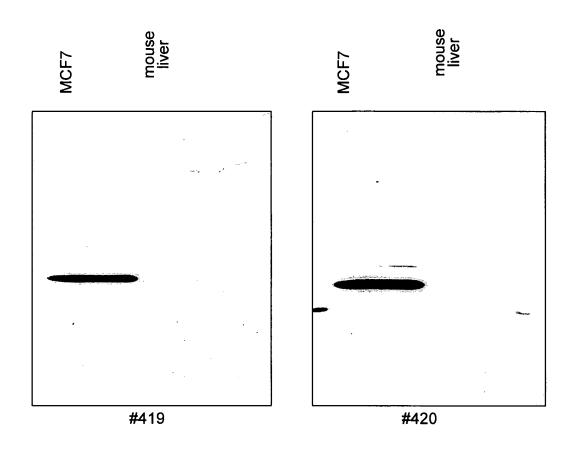


Figure 5B

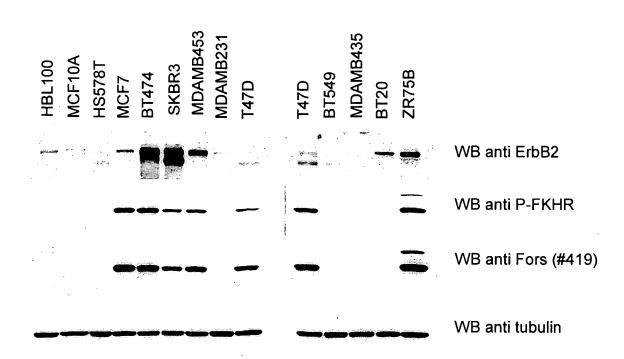
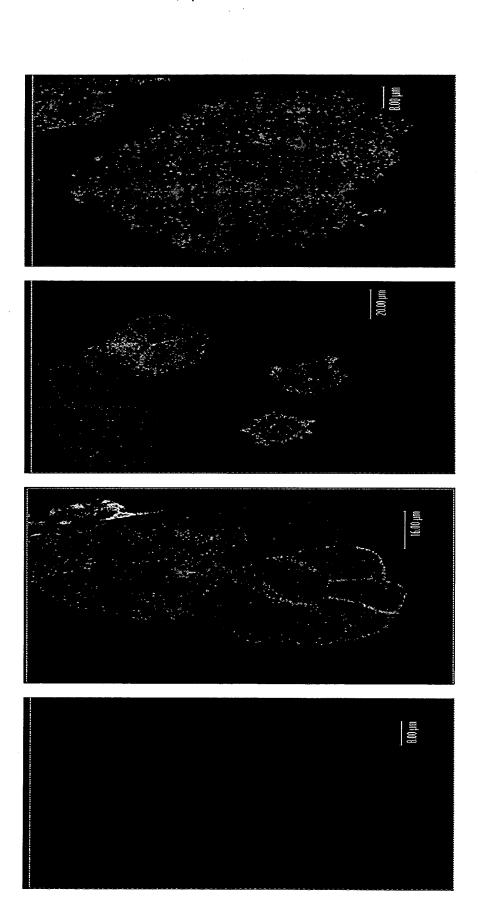


Figure 5C





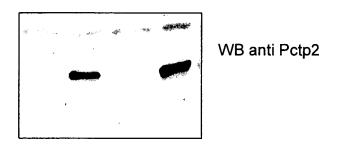


Figure 6A

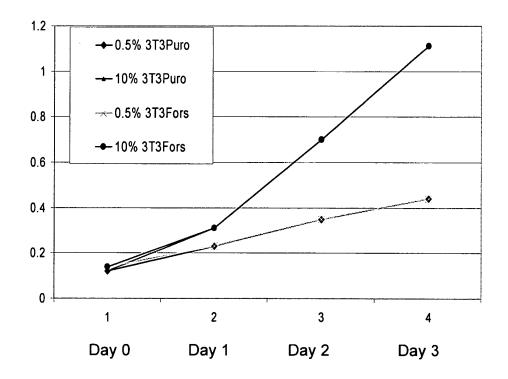


Figure 6B

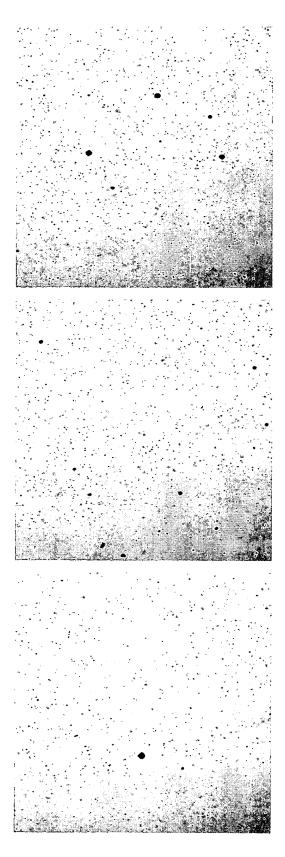
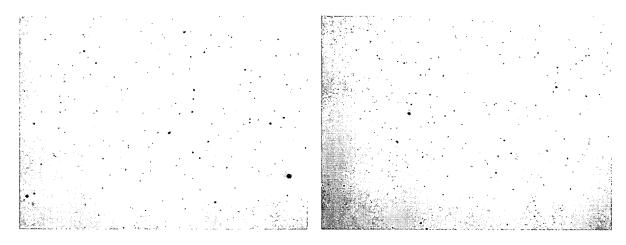


Figure 7A

HER14 Puro



HER14 Fors

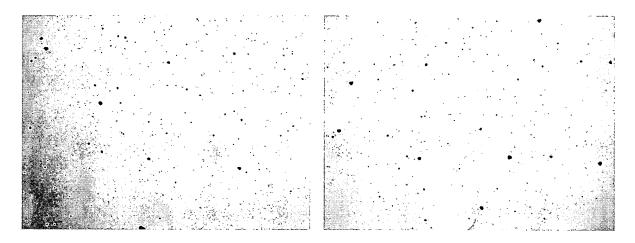
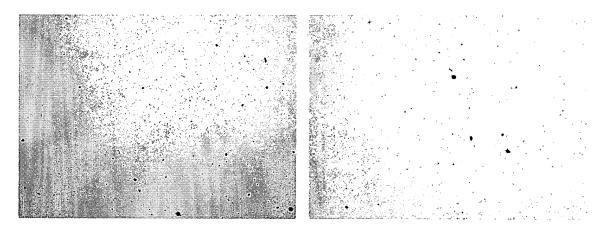


Figure 7B

HER14 Puro + 10 ng/ml EGF



HER14 Fors + 10 ng/ml EGF

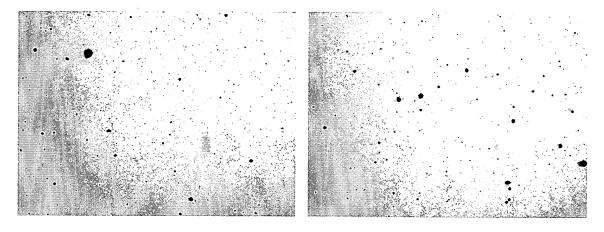


Figure 7C

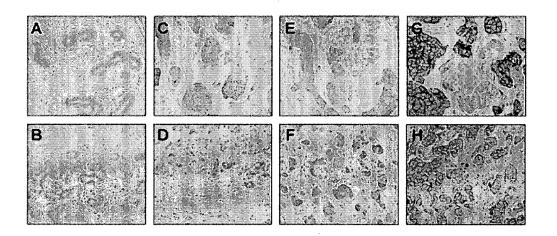
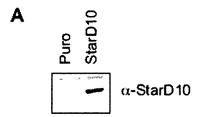


Figure 8 A-H



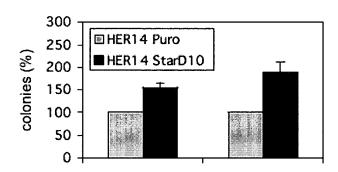


Figure 9



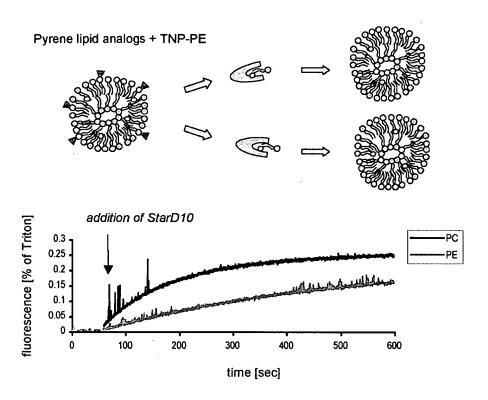


Figure 10

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